

More about ANOVA

An F-test in an analysis of variance is actually a test for a specific comparison of two two hypothesis, each specifying a model.

In the one-way ANOVA,

$$H_0: \mu_1 = \mu_2 = \dots = \mu_g = \mu^*$$

$$\text{or } \alpha_1 = \alpha_2 = \dots \alpha_g = 0$$

$$\text{Model is } y_{ij} = \mu^* + \epsilon_{ij}$$

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H_a : At least two μ_i 's differ
or at least two α_i 's differ
Model is $y_{ij} = \mu_i + \epsilon_{ij} = \mu^* + \alpha_i + \epsilon_{ij}$
As a model, H_a is sometimes called the *unrestricted model* or the *full model*.

Suppose you knew H_0 were true.

- Your best estimate of μ^* would be $\bar{y}_{..}$
- The residuals would be $y_{ij} - \bar{y}_{..}$
- The residual SS would be $SSR_0 = SS_\tau = \sum \sum (y_{ij} - \bar{y}_{..})^2$.

In the unrestricted case (H_A),

- Your best estimates of μ_1, \dots, μ_g are $\bar{y}_{1.}, \dots, \bar{y}_{g.}$
- The residuals would be $y_{ij} - \bar{y}_{i.}$
- The residual SS would be $SSR_A = SS_E = \sum_i \sum_j (y_{ij} - \bar{y}_{i.})^2$.

Thus $SS_{trt} = SS_\tau - SS_E = SSR_0 - SSR_A$ is the *reduction in the residual SS* you can achieve if you leave H_0 in favor of H_a and

$$F = \{SS_{trt} / (g-1)\} / \{SS_E / (N-g)\}$$

is a way to see if this reduction is large enough to be significant.

This is a general principle used in ANOVA and regression:

$$F = \frac{(SSR_0 - SSR_A) / (df_0 - df_A)}{SSR_A / df_A}$$

where

$$df_A = N - n_A, n_A = \# \text{parameters for } H_a) \\ = N - g \quad (\text{in this case})$$

and

$$df_0 = N - n_0, n_0 = \# \text{parameters for } H_0) \\ = N - 1 \quad (\text{in this case}) \\ df_{trt} = df_0 - df_A = N - 1 - (N - g) = g - 1 \\ df_{error} = df_A = N - g$$

Comment The ratio SS_{trt} / SS_τ is the proportion of the total variation that can be “explained” by differential effects of treatments. It is the direct analogue of the coefficient of determination (multiple R^2) in regression.

Why is this effective? It all depends on the **expectations of mean squares** (MS) in the ANOVA.

Suppose H_0 is true. Then

$$E[SSR_0] = df_0 \sigma^2 = (N - 1) \sigma^2$$

and

$$E[SSR_A] = \sum (n_i - 1) \sigma^2 = df_A \sigma^2 = (N - g) \sigma^2$$

Therefore

$$E[SS_{trt}] = E[SSR_0] - E[SSR_A]$$

$$(N - 1) \sigma^2 - (N - g) \sigma^2 = (g - 1) \sigma^2 = df_{trt} \sigma^2$$

$$E[SS_{error}] = E[SSR_1] = (N - g) \sigma^2 = df_{error} \sigma^2$$

Since mean squares are SS/df ,

$$E[MS_{trt}] = E[SS_{trt} / (g - 1)] = \sigma^2$$

$$E[MS_{error}] = E[SS_{error} / (N - g)] = \sigma^2$$

Conclusion: When H_0 is true, the expectation of both the numerator and denominator of $F = MS_{trt} / MS_{error}$ are the same. The median of F is close to 1.

When H_0 is not true, it is still true that

$$E[MS_{error}] = E[SS_{error} / (N - g)] = \sigma^2$$

Now, however,

$$E[MS_{trt}] = \sigma^2 + \tau^2 / (g - 1) > \sigma^2$$

where

$$\tau^2 \equiv \sum_{1 \leq i \leq g} n_i (\mu_i - \tilde{\mu})^2, \quad \tilde{\mu} = \sum n_i \mu_i / N.$$

Note that $\tau^2 = 0$ when H_0 is true.

So violation of H_0 increases $E[MS_{trt}]$ and hence $E[F]$, and makes it more probable you will reject H_0 .

If you use the parametrization which sets

$$\mu^* = \tilde{\mu} = \sum n_i \mu_i / N \text{ and } \alpha_i = \mu_i - \mu^*,$$

$$\tau^2 \equiv \sum_{1 \leq i \leq g} n_i \alpha_i^2$$

This formula is *not* correct for other choices for μ^* . In particular it is not true for $\mu^* = \bar{\mu} = \sum \mu_i / g$, $\alpha_i = \mu_i - \bar{\mu}$ (unless the sample sizes are equal).

To summarize:

Testing an ANOVA hypothesis is equivalent to a comparison of two models

- a null model
- and

- a more general alternative model.

Your conclusion in the test is effectively a *selection* of one or the other model as most appropriate.

In more complex ANOVA's you may be selecting among more than two models.

Contrasts

A contrast is a formula which compares two or more treatment means or effects in a way that doesn't depend on the overall level μ^* .

Examples:

- $\mu_1 - \mu_3 = (\mu^* + \alpha_1) - (\mu^* + \alpha_3) = \alpha_1 - \alpha_3$
- $(\mu_1 + \mu_2)/2 - (\mu_3 + \mu_4 + \mu_5)/3$
 $= (\mu^* + \alpha_1 + \mu^* + \alpha_2)/2$
 $- (\mu^* + \alpha_3 + \mu^* + \alpha_4 + \mu^* + \alpha_5)/3$
 $= (\alpha_1 + \alpha_2)/2 - (\alpha_3 + \alpha_4 + \alpha_5)/3$

This compares the average of the first 2 means or effects with the average of the last 3.

Formal definition

A *contrast* is a linear combination of μ 's

$$w(\{\mu_i\}) \equiv \sum_i w_i \mu_i, \text{ with } \sum_i w_i = 0$$

Because $\sum_i w_i = 0$, $w(\{\mu_i\})$ doesn't depend on μ^* :

$$\begin{aligned} \sum_i w_i \mu_i &= \sum_i w_i (\mu^* + \alpha_i) \\ &= (\sum_i w_i) \mu^* + \sum_i w_i \alpha_i = \sum_i w_i \alpha_i \\ &= 0 \times \mu^* + \sum_i w_i \alpha_i = \sum_i w_i \alpha_i \\ &= w(\{\alpha_i\}) \end{aligned}$$

Since $\sum_i w_i \alpha_i$ doesn't depend on μ^* this satisfies the informal definition of a contrast given before.

The weights $\{w_i\}$ themselves are also often referred to as a *contrast*.

An *observed contrast* is

$$w(\{\overline{y}_i\}) = \sum_i w_i \overline{y}_i = \sum_i w_i \hat{\alpha}_i = w(\{\hat{\alpha}_i\})$$

You may sometime calculate several contrasts as part of your analysis. What you use depends on the *questions of interest* to the researcher, not on some statistical magic.

If you are just providing statistical advice, you need to find out what questions need answers.

More on the example:

```
Cmd> anova("Logy=treat",fstat:T)
Model used is logy = treat
WARNING: summaries are sequential
```

	DF	SS	MS	F	P-value
CONSTANT	1	79.425	79.425	8653.95365	< 1e-08
treat	4	3.5376	0.88441	96.36296	< 1e-08
ERROR1	32	0.29369	0.0091779		

Compute $\hat{\mu}_i = \overline{y}_i$, and $\hat{\alpha}_i = \overline{y}_i - \sum \overline{y}_i / g$

```
Cmd> muhats <- tabs(logy,treat,mean:T);muhats
(1) 1.9325 1.6287 1.3775 1.1943 1.0567
Cmd> alphahats <- muhats - sum(muhats)/5; alphahats
(1) 0.49456 0.19081 -0.06044 -0.24365 -0.38127
```

MacAnova function `coefs()` computes $\hat{\alpha}_i$'s

```
Cmd> coefs(treat) # or coefs("treat") or coefs(2)
(1) 0.49456 0.19081 -0.06044 -0.24365 -0.38127
treat is line 2 in anova() output.
```

Enter weights and compute contrast two ways.

```
Cmd> w <- vector(vector(1,1)/2,-vector(1,1,1)/3); w
(1) 0.5 0.5 -0.33333 -0.33333 -0.33333
Cmd> vector(sum(w),sum(w*muhats),sum(w*alphahats))
(1) 1.1102e-16 0.57114 0.57114
```

MacAnova function `contrast()` makes it easy to compute contrasts.

```
Cmd> stuff <- contrast(treat, w); stuff
component: estimate      Same as just computed
(1) 0.57114
component: ss
(1) 2.9446
component: se
(1) 0.031886
```

The result (output) from `contrast()` is a *structure* with three components:

- The estimate component is the value of the contrast $\sum W_i \alpha_i$.
- The se component is its estimated standard error. You can compute a t-statistic to test the null hypothesis that the $\sum W_i \alpha_i = 0$
- The ss component is an SS associated with the contrast.

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Using estimate and se to compute a t-statistic to test $H_0: \sum W_i \alpha_i = 0$:

```
Cmd> tstat <- stuff$estimate/stuff$se
Cmd> vector(tstat,twotailt(tstat,DF[3]))
(1) 17.912 3.0663e-18 t-statistic and p-value
```

When H_0 is true, t has Student's t-distribution on $df_{\text{error}} = N - g$ d.f.

`stuff$estimate` is one way to extract a component from a structure. Since this is the first component, another way is `stuff[[1]]` and it is `stuff[[1]]/stuff[[3]]`.

The ss component (`stuff$ss` or `stuff[[2]]`) is $MSE_{\text{estimate}}^2 / se^2$

```
Cmd> mse <- SS[3]/DF[3]; mse # MS in ERROR1 row of ANOVA
      ERROR1
0.0091779
Cmd> mse*(stuff$estimate/stuff$se)^2
(1) 2.9446
Cmd> vector(stuff$ss/mse, tstat^2)
(1) 320.83 320.83
```

`stuff$ss/mse` is the same as t^2 :

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Common Contrasts

- Pairwise contrasts

$$\mu_i - \mu_j = \alpha_i - \alpha_j \quad \text{Compare two groups}$$

$$\{w_{ij}\} = \{0, \dots, 1, 0, \dots, -1, 0, \dots\}$$

For g groups, there are $g(g-1)/2$ essentially different pairwise contrasts:

```
Cmd> print(pairwise_wts, format="4.0F")
pairwise_wts: Each column is a set of contrast weights
(1,1) 1 1 1 1 0 0 0 0 0 0
(2,1) -1 0 0 0 1 1 1 0 0 0
(3,1) 0 -1 0 0 -1 0 0 1 1 0
(4,1) 0 0 0 -1 0 0 -1 0 1 1
(5,1) 0 0 0 0 -1 0 0 0 -1 -1
```

The following computes the contrast and t-statistic for the $5 \times 4/2 = 10$ pairwise contrasts.

```
Cmd> for(i,1,10){
  stuff <- contrast(treat,pairwise_wts[,i])
  print(paste("W =", pairwise_wts[,i], "estimate =", \
    stuff$estimate, " t-statistic =", \
    stuff$estimate/stuff$se))
}
W = 1 -1 0 0 0 , estimate = 0.30375 , t-statistic = 6.3413
W = 1 0 -1 0 0 , estimate = 0.555 , t-statistic = 11.586
W = 1 0 0 -1 0 , estimate = 0.73821 , t-statistic = 14.889
W = 1 0 0 0 -1 , estimate = 0.87583 , t-statistic = 16.928
W = 0 1 -1 0 0 , estimate = 0.25125 , t-statistic = 5.2452
W = 0 1 0 -1 0 , estimate = 0.43446 , t-statistic = 8.7626
W = 0 1 0 0 -1 , estimate = 0.57208 , t-statistic = 11.057
W = 0 0 1 -1 0 , estimate = 0.18321 , t-statistic = 3.6952
W = 0 0 1 0 -1 , estimate = 0.32083 , t-statistic = 6.201
W = 0 0 0 1 -1 , estimate = 0.13762 , t-statistic = 2.582
```

- Comparison with control

Say treatment 1 is a control.

An obvious idea is to compare the mean or effect of the control with the average mean or effect of all the non-controls:

$$\mu_1 - (\mu_2 + \mu_3 + \dots + \mu_g)/(g-1)$$

$$= \alpha_1 - (\alpha_2 + \alpha_3 + \dots + \alpha_g)/(g-1)$$

Contrast coefficients are

$$\{w_{ij}\} = \{1, -1/(g-1), \dots, -1/(g-1)\}$$

Of course individual pairwise comparisons with control $\alpha_i - \alpha_1$, $i = 2, \dots, g$ would probably of interest too.

Multiplying this by $g - 1$, an equivalent contrast is

$$(g-1)\mu_1 - \mu_2 - \mu_3 - \dots - \mu_g$$

with integer coefficients $\{g-1, -1, \dots, -1\}$. Before computers were common, this made calculations easier.

- **Factorial treatments**

When there are two factors, A and B, each at two levels, there are 4 treatments with means μ_{11} , μ_{12} , μ_{21} , μ_{22} . These can be displayed in a 2 by 2 table

	B ₁	B ₂
A ₁	μ_{11}	μ_{12}
A ₂	μ_{21}	μ_{22}

Natural contrasts would be

- Average of row 1 vs average of row 2:
 $(\mu_{11} + \mu_{12})/2 - (\mu_{21} + \mu_{22})/2$

This measures the effect of factor A, ignoring factor B (main effect of A).

- Average of col. 1 vs average of col. 2:
 $(\mu_{11} + \mu_{21})/2 - (\mu_{12} + \mu_{22})/2$

This measures the effect of factor B, ignoring factor A (main effect of B).

- Difference between effects of A for the two levels of B

$$(\mu_{11} - \mu_{21}) - (\mu_{12} - \mu_{22})$$

This is algebraically the same as the difference between effects of B for the two levels of A

$$(\mu_{11} - \mu_{12}) - (\mu_{21} - \mu_{22})$$

When this contrast is not zero, it means the effect of A depends on the level of B (or the effect of B depends on the level of A).

When this occurs, we say there is *interaction* between factors A and B. So this is an *interaction* contrast.

Suppose the treatments are determined by a quantitative variable x with levels x_1, x_2, \dots, x_g , say. Then, if you fit a straight line, the least squares estimate of the slope is

$$b = \frac{\sum n_i(x_i - \bar{x})y_i}{\sum n_i(x_i - \bar{x})^2}, \quad \bar{x} = \frac{\sum n_i x_i}{N}$$

When the sample sizes are equal, you can omit the n_i .

This is a contrast with weights

$$w_i = n_i(x_i - \bar{x}) / \sum_i n_i(x_i - \bar{x})^2$$

which do satisfy $\sum w_i = 0$, because $\sum_i n_i(x_i - \bar{x}) = 0$.

It will be large when there is a high degree of linear dependence of the means on x .

This is a *linear* contrast because it focuses on the strength of a straight line relationship between μ_i or α_i and x_i .